SEQUENCE LISTING

<110>	110> Zhou, Qun-Yong Ehlert, Frederick														
	:120> Prokineticin Polypeptides, Related Compositions and Methods														
<130>	:130> P-UC 5016														
	:150> 60/245,882 :151> 2000-11-03														
<160>	<160> 19														
<170>	170> FastSEQ for Windows Version 4.0														
<211> <212>	2210> 1 2211> 1377 2212> DNA 2213> Homo sapiens														
<220> <221> CDS <222> (55)(369)															
	400> 1 gggaagega gaggeateta ageaggeagt gttttgeett caccecaagt gace													atg Met 1	57
aga g Arg G	gt gco ly Ala	_	_	-			_								105
gac t Asp C		a Val				_	-			_	-				153
Ala G	gc aco ly Thi 35	_	_	-		_	_			_				_	201
_	cc cco hr Pro	_			_				_				_		249
	tc cco al Pro														297
ccc a	ac cts	g ctg	tgc	tcc	agg	ttc	ccg	gac	ggc	agg	tac	cgc	tgc	tcc	345

```
Pro Asn Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys Ser 85 90 95
```

atg gac ttg aag aac atc aat ttt taggcgcttg cctggtctca ggatacccac 399
Met Asp Leu Lys Asn Ile Asn Phe
100 105

catcetttte tgageacage etggattttt atttetgeca tgaaacccag eteccatgae 459 teteceagte cetacactga etaceetgat etetettgte tagtacgeae atatgeacae 519 aggcagacat acctcccatc atgacatggt ccccaggctg gcctgaggat gtcacagctt 579 gaggetgtgg tgtgaaaggt ggecageetg gttetettee etgeteagge tgecagagag 639 gtggtaaatg gcagaaagga cattccccct ccctcccca ggtgacctgc tctctttcct 699 gggccctgcc cctctcccca catgtatccc tcqqtctqaa ttaqacattc ctqqqcacaq 759 getettgggt geattgetea gagteceagg teetggeetg acceteagge cetteaegtg 819 aggtetgtga ggaccaattt gtgggtagtt catetteect egattggtta acteettagt 879 ttcagaccac agactcaaga ttggctcttc ccagagggca gcagacagtc accccaaggc 939 aggtgtaggg agcccaggga ggccaatcag cccctgaag actctggtcc cagtcagcct 999 gtggcttgtg gcctgtgacc tqtgaccttc tqccaqaatt gtcatqcctc tqagqcccc 1059 tettaceaca etttaceagt taaceaetga ageeeccaat teecacaget tttecattaa 1119 aatgcaaatg gtggtggttc aatctaatct gatattgaca tattagaagg caattagggt 1179 gtttccttaa acaactcctt tccaaggatc agccctgaga gcaggttggt gactttgagg 1239 agggcagtcc tctgtccaga ttggggtggg agcaagggac agggagcagg gcaggggctg 1299 aaaggggcac tgattcagac cagggaggca actacacacc aacctgctgg ctttagaata 1359 aaagcaccaa ctgaactg 1377

<210> 2 <211> 105 <212> PRT

<213> Homo sapiens

<400> 2

 Met Arg Gly Ala Thr Arg Val Ser Ile Met Leu Leu Leu Leu Val Thr Val

 1
 5
 10
 15

 Ser Asp Cys Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys
 20
 25
 30

 Gly Ala Gly Thr Cys Cys Ala Ile Ser Leu Trp Leu Arg Gly Leu Arg
 35
 40
 45

45 Thr Pro Leu Glv Ara Glu Glv Glu Glu Cvs His

Met Cys Thr Pro Leu Gly Arg Glu Gly Glu Glu Cys His Pro Gly Ser 50 55 60

His Lys Val Pro Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys 70 75 80

Leu Pro Asn Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys
85 90 95

Ser Met Asp Leu Lys Asn Ile Asn Phe 100 105

<210> 3

<211> 86

<212> PRT

<213> Homo sapiens

<400> 3

Ala 1	Val	Ile	Thr	Gly 5	Ala	Cys	Glu	Arg	Asp	Val	Gln	Cys	Gly	Ala 15	Gly
Thr	Cys	Cys	Ala 20	Ile	Ser	Leu	Trp	Leu 25	Arg	Gly	Leu	Arg	Met 30	Cys	Thr
Pro	Leu	Gly 35	Arg	Glu	Gly	Glu	Glu 40	Cys	His	Pro	Gly	Ser 45	His	Lys	Val
Pro	Phe 50	Phe	Arg	Lys	Arg	Lys 55	His	His	Thr	Cys	Pro 60	Cys	Leu	Pro	Asn
Leu 65	Leu	Cys	Ser	Arg	Phe 70	Pro	Asp	Gly	Arg	Tyr 75	Arg	Cys	Ser	Met	Asp 80
Leu	Lys	Asn	Ile	Asn 85	Phe										

<210> 4 <211> 1406 <212> DNA

<213> Homo sapiens

<220> <221> CDS <222> (10)...(333)

<400> 4

gagggegee atg agg age etg tge tge gee eea etc etg etc etc ttg etg 51 Met Arg Ser Leu Cys Cys Ala Pro Leu Leu Leu Leu Leu Leu 1 5 10

ctg ccg ccg ctg ctc acg ccc cgc gct ggg gac gcc gcc gtg atc 99
Leu Pro Pro Leu Leu Thr Pro Arg Ala Gly Asp Ala Ala Val Ile
15 20 25 30

acc ggg gct tgt gac aag gac tcc caa tgt ggt gga ggc atg tgc tgt 147
Thr Gly Ala Cys Asp Lys Asp Ser Gln Cys Gly Gly Met Cys Cys
35 40 45

gct gtc agt atc tgg gtc aag agc ata agg att tgc aca cct atg ggc 195
Ala Val Ser Ile Trp Val Lys Ser Ile Arg Ile Cys Thr Pro Met Gly
50 55 60

aaa ctg gga gac agc tgc cat cca ctg act cgt aaa gtt cca ttt ttt 243 Lys Leu Gly Asp Ser Cys His Pro Leu Thr Arg Lys Val Pro Phe Phe 65 70 75

ggg cgg agg atg cat cac act tgc cca tgt ctg cca ggc ttg gcc tgt 291
Gly Arg Arg Met His His Thr Cys Pro Cys Leu Pro Gly Leu Ala Cys
80 85 90

tta cgg act tca ttt aac cga ttt att tgt tta gcc caa aag 333 Leu Arg Thr Ser Phe Asn Arg Phe Ile Cys Leu Ala Gln Lys 95 100 105

taatcgctct ggagtagaaa ccaaatgtga atagccacat cttacctgta aagtcttact 393

```
tgtgattgtg ccaaacaaaa aatgtgccag aaagaaatgc tcttgcttcc tcaactttcc 453
aagtaacatt tttatetttg atttgtaaat gatttttttt tttttttta tegaaagaga 513
attttacttt tggatagaaa tatgaagtgt aaggcattat ggaactggtt cttatttccc 573
tgtttgtgtt ttggtttgat ttggcttttt tcttaaatgt caaaaacgta cccattttca 633
caaaaatgag gaaaataaga atttgatatt ttgttagaaa aactttttt ttttttctc 693
accaccccaa geceeatttg tgeeetgeeg cacaaataca ectacagett ttggteeett 753
ttccctcctc ttgcatttta aagtggaggg tttgtctctt tgagtttqat qqcaqaatca 873
ctgatgggaa tccagctttt tgctggcatt taaatagtga aaaqagtgta tatqtqaact 933
tgacactcca aactcctgtc atggcacgga agctaggagt gctgctggac ccttcctaaa 993
cctgtcactc aagaggactt cagctctgct gttgggctgg tgtgtggaca gaaggaatgg 1053
aaagccaaat taatttagtc cagatttcta ggtttgggtt tttctaaaaa taaaagatta 1113
catttacttc ttttactttt tataaagttt tttttcctta gtctcctact tagagatatt 1173
ctagaaaatg tcacttgaag aggaagtatt tattttaatc tqqcacaaca ctaattacca 1233
tttttaaagc ggtattaagt tgtaatttaa accttgtttg taactgaaag gtcgattgta 1293
atggattgcc gtttgtacct gtatcagtat tgctgtgtaa aaattctgta tcagaataat 1353
aacagtactg tatatcattt gatttatttt aatattatat ccttattttt qtc
<210> 5
<211> 108
<212> PRT
<213> Homo sapiens
<400> 5
Met Arg Ser Leu Cys Cys Ala Pro Leu Leu Leu Leu Leu Leu Pro
Pro Leu Leu Thr Pro Arg Ala Gly Asp Ala Ala Val Ile Thr Gly
Ala Cys Asp Lys Asp Ser Gln Cys Gly Gly Met Cys Cys Ala Val
                           40
Ser Ile Trp Val Lys Ser Ile Arg Ile Cys Thr Pro Met Gly Lys Leu
                       55
Gly Asp Ser Cys His Pro Leu Thr Arg Lys Val Pro Phe Phe Gly Arg
                   70
                                      75
Arg Met His His Thr Cys Pro Cys Leu Pro Gly Leu Ala Cys Leu Arg
Thr Ser Phe Asn Arg Phe Ile Cys Leu Ala Gln Lys
           100
                               105
<210> 6
<211> 81
<212> PRT
<213> Homo sapiens
<400> 6
Ala Val Ile Thr Gly Ala Cys Asp Lys Asp Ser Gln Cys Gly Gly
                                  10
Met Cys Cys Ala Val Ser Ile Trp Val Lys Ser Ile Arg Ile Cys Thr
                               25
Pro Met Gly Lys Leu Gly Asp Ser Cys His Pro Leu Thr Arg Lys Val
```

35 40 45
Pro Phe Phe Gly Arg Arg Met His His Thr Cys Pro Cys Leu Pro Gly

```
50
                        55
                                            60
Leu Ala Cys Leu Arg Thr Ser Phe Asn Arg Phe Ile Cys Leu Ala Gln
                    70
                                        75
Lys
<210> 7
<211> 21
<212> PRT
<213> Homo sapiens
<400> 7
Asn Asn Phe Gly Asn Gly Arg Gln Glu Arg Arg Lys Arg Lys Arg Ser
                                    10
Lys Arg Lys Lys Glu
<210> 8
<211> 21
<212> PRT
<213> Homo sapiens
<400> 8
Ser His Val Ala Asn Gly Arg Gln Glu Arg Arg Arg Ala Lys Arg Arg
                                    10
Lys Arg Lys Lys Glu
            20
<210> 9
<211> 19
<212> PRT
<213> Homo sapiens
<400> 9
Met Arg Gly Ala Thr Arg Val Ser Ile Met Leu Leu Val Thr Val
Ser Asp Cys
<210> 10
<211> 26
<212> PRT
<213> Homo sapiens
<400> 10
Met Arg Ser Leu Cys Cys Ala Pro Leu Leu Leu Leu Leu Leu Pro
Leu Leu Thr Pro Pro Ala Gly Asp Ala
```

```
<210> 11
<211> 96
<212> PRT
<213> Bombina variegata
<400> 11
Met Lys Cys Phe Ala Gln Ile Val Val Leu Leu Val Ile Ala Phe
                                    10
Ser His Gly Ala Val Ile Thr Gly Ala Cys Asp Lys Asp Val Gln Cys
Gly Ser Gly Thr Cys Cys Ala Ala Ser Ala Trp Ser Arg Asn Ile Arg
Phe Cys Ile Pro Leu Gly Asn Ser Gly Glu Asp Cys His Pro Ala Ser
His Lys Val Pro Tyr Asp Gly Lys Arg Leu Ser Ser Leu Cys Pro Cys
                    70
                                        75
Lys Ser Gly Leu Thr Cys Ser Lys Ser Gly Glu Lys Phe Lys Cys Ser
                                    90
<210> 12
<211> 81
<212> PRT
<213> Dendroaspis polylepis polylepis
<400> 12
Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Leu Gln Cys Gly Lys Gly
Thr Cys Cys Ala Val Ser Leu Trp Ile Lys Ser Val Arg Val Cys Thr
Pro Val Gly Thr Ser Gly Glu Asp Cys His Pro Ala Ser His Lys Ile
                            40
Pro Phe Ser Gly Gln Arg Lys Met His His Thr Cys Pro Cys Ala Pro
Asn Leu Ala Cys Val Gln Thr Ser Pro Lys Lys Phe Lys Cys Leu Ser
65
                    70
Lys
<210> 13
<211> 81
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 13
Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys Gly Ala Gly
```

<210> 14

<211> 86

<212> PRT

<213> Artificial Sequence

<220>

<223> synthetic construct

<400> 14

Ala Val Ile Thr Gly Ala Cys Asp Lys Asp Ser Gln Cys Gly Gly 1 5 10 15

Met Cys Cys Ala Val Ser Ile Trp Val Lys Ser Ile Arg Ile Cys Thr
20 25 30

Pro Met Gly Lys Leu Gly Asp Ser Cys His Pro Leu Thr Arg Lys Val 35 40 45

Pro Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys Leu Pro Asn 50 55 60

Leu Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys Ser Met Asp 65 70 75 80

Leu Lys Asn Ile Asn Phe

85

<210> 15

<211> 89

<212> PRT

<213> Artificial Sequence

<220>

<223> synthetic construct

<400> 15

Gly Ile Leu Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys
1 10 15

Gly Ala Gly Thr Cys Cys Ala Ile Ser Leu Trp Leu Arg Gly Leu Arg
20 25 30

Met Cys Thr Pro Leu Gly Arg Glu Gly Glu Glu Cys His Pro Gly Ser 35 40 45

His Lys Val Pro Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys 50 55 60

Leu Pro Asn Leu Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys

```
65
                    70
                                        75
                                                            80
Ser Met Asp Leu Lys Asn Ile Asn Phe
                85
<210> 16
<211> 85
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 16
Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys Gly Ala Gly Thr
Cys Cys Ala Ile Ser Leu Trp Leu Arg Gly Leu Arg Met Cys Thr Pro
                                25
Leu Gly Arg Glu Gly Glu Cys His Pro Gly Ser His Lys Val Pro
Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys Leu Pro Asn Leu
Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys Ser Met Asp Leu
Lys Asn Ile Asn Phe
<210> 17
<211> 86
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 17
Ala Ala Ala Ala Ala Cys Glu Arg Asp Val Gln Cys Gly Ala Gly
Thr Cys Cys Ala Ile Ser Leu Trp Leu Arg Gly Leu Arg Met Cys Thr
Pro Leu Gly Arg Glu Gly Glu Cys His Pro Gly Ser His Lys Val
Pro Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys Leu Pro Asn
Leu Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys Ser Met Asp
```

<210> 18 <211> 87

Leu Lys Asn Ile Asn Phe

85

```
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 18
Met Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys Gly Ala
Gly Thr Cys Cys Ala Ile Ser Leu Trp Leu Arg Gly Leu Arg Met Cys
                                25
Thr Pro Leu Gly Arg Glu Gly Glu Cys His Pro Gly Ser His Lys
Val Pro Phe Phe Arg Lys Arg Lys His His Thr Cys Pro Cys Leu Pro
                        55
Asn Leu Cys Ser Arg Phe Pro Asp Gly Arg Tyr Arg Cys Ser Met
Asp Leu Lys Asn Ile Asn Phe
                85
<210> 19
<211> 14
<212> PRT
<213> Artificial Sequence
<220>
<223> synthetic construct
<400> 19
Ala Val Ile Thr Gly Ala Cys Glu Arg Asp Val Gln Cys Gly
```